

CD103PCT.ST25.txt
SEQUENCE LISTING

<110> CropDesign N.V.

<120> Plants having modified growth characteristics and method for making the same

<130> CD-103-PCT

<150> EP 03077811.2

<151> 2003-09-05

<160> 21

<170> PatentIn version 3.3

<210> 1

<211> 930

<212> DNA

<213> Arabidopsis thaliana

```

<400> 1
atggagaagt acgagaagct agagaaggtc ggagaaggaa catacgggaa agtctacaaa 60
gcgatggaga aaggaactgg taagcttggt gctctgaaga aaactcgtct cgagatggac 120
gaagaaggta ttccaccaac tgctcttcgt gagatctcgc ttctccagat gttatcaaca 180
tcgatctatg ttgttcgatt actctgcgtc gaacatgttc atcaaccatc aaccaaattc 240
caatctacca aatccaatct ctatctcggt ttcgagtatc tcgatactga tcttaagaaa 300
ttcatcgatt cgtataggaa aggacctaat cctaagcctc ttgagccttt tttgattcag 360
aagttgatgt ttcagctttg taaagggtgt gcgcattgtc atagtcattg tgtgcttcac 420
cgtgatctta aaccgcagaa tcttctctcg gtgaaagata aagagcttct taagattgct 480
gatttgggtc ttggtcgtgc ttttactggt cctcttaagt cttatacgca tgagattggt 540
actctttggt atagagctcc tgaagttcct cttggatcta ctcattattc aactggtgtt 600
gacatgtggt ctggttggtg tatctttgct gagatgggtc ggaggcaagc tcttttccct 660
ggtgattctg agtttcagca attgcttcat atcttcaggt tgctaggaac accaactgag 720
cagcaatggc cgggtgtttc cacactgcgt gactggcatg tttaccctaa gtgggagccg 780
caagacttaa ctcttgctgt tccttctctt tcacctcaag gagttgatct tctcacgaaa 840
atgctcaagt acaatccagc cgaaagaatt tcagcaaaaa cagcacttga tcaccatata 900
tttgacagcc ttgacaagtc tcagttctga 930

```

<210> 2

<211> 309

<212> PRT

<213> Arabidopsis thaliana

<400> 2

```

Met Glu Lys Tyr Glu Lys Leu Glu Lys Val Gly Glu Gly Thr Tyr Gly
1 5 10 15

```

```

Lys Val Tyr Lys Ala Met Glu Lys Gly Thr Gly Lys Leu Val Ala Leu
20 25 30

```

```

Lys Lys Thr Arg Leu Glu Met Asp Glu Glu Gly Ile Pro Pro Thr Ala
35 40 45

```

```

Leu Arg Glu Ile Ser Leu Leu Gln Met Leu Ser Thr Ser Ile Tyr Val
50 55 60

```

```

Val Arg Leu Leu Cys Val Glu His Val His Gln Pro Ser Thr Lys Ser
65 70 75 80

```

CD103PCT.ST25.txt

Gln Ser Thr Lys Ser Asn Leu Tyr Leu Val Phe Glu Tyr Leu Asp Thr
 85 90 95
 Asp Leu Lys Lys Phe Ile Asp Ser Tyr Arg Lys Gly Pro Asn Pro Lys
 100 105 110
 Pro Leu Glu Pro Phe Leu Ile Gln Lys Leu Met Phe Gln Leu Cys Lys
 115 120 125
 Gly Val Ala His Cys His Ser His Gly Val Leu His Arg Asp Leu Lys
 130 135 140
 Pro Gln Asn Leu Leu Leu Val Lys Asp Lys Glu Leu Leu Lys Ile Ala
 145 150 155 160
 Asp Leu Gly Leu Gly Arg Ala Phe Thr Val Pro Leu Lys Ser Tyr Thr
 165 170 175
 His Glu Ile Val Thr Leu Trp Tyr Arg Ala Pro Glu Val Leu Leu Gly
 180 185 190
 Ser Thr His Tyr Ser Thr Gly Val Asp Met Trp Ser Val Gly Cys Ile
 195 200 205
 Phe Ala Glu Met Val Arg Arg Gln Ala Leu Phe Pro Gly Asp Ser Glu
 210 215 220
 Phe Gln Gln Leu Leu His Ile Phe Arg Leu Leu Gly Thr Pro Thr Glu
 225 230 235 240
 Gln Gln Trp Pro Gly Val Ser Thr Leu Arg Asp Trp His Val Tyr Pro
 245 250 255
 Lys Trp Glu Pro Gln Asp Leu Thr Leu Ala Val Pro Ser Leu Ser Pro
 260 265 270
 Gln Gly Val Asp Leu Leu Thr Lys Met Leu Lys Tyr Asn Pro Ala Glu
 275 280 285
 Arg Ile Ser Ala Lys Thr Ala Leu Asp His Pro Tyr Phe Asp Ser Leu
 290 295 300
 Asp Lys Ser Gln Phe
 305
 <210> 3
 <211> 936
 <212> DNA
 <213> Arabidopsis thaliana
 <400> 3
 atggagaaat acgagaagct cgaaaaggct ggtgaaggaa cctatggaaa agtctacaaa 60
 gcaatggaga aaaccaccgg aaaactcgtc gctctgaaga aaactaggct cgaaatggac 120
 gaagaaggta taccaccaac ggctctccgt gagatctctc ttctccaaat gctttctcaa 180
 tcaatctaca tcgttcgtct cctctgcgtc gaacatgtta ttcaatcgaa agattcgact 240
 gtttctcact ctcccaaadc caatctctat ctcgtttttg agtatctcga cactgatctc 300
 aagaaattta tagattctca tagaaagggc tcgaatccta gaccgcttga ggcttctctt 360
 gtgcagaggt ttatgtttca gctttttaaa ggtgtggctc attgtcatag ccatgggtgtg 420

CD103PCT.ST25.txt

```

cttcaccgtg atcttaaacc gcagaatctt ctattggata aggataaagg gattcttaag 480
attgctgatt tgggtcttag tcgtgctttt actgtgcctc ttaaggctta tacacatgag 540
attgttactc tttggtatag agctcctgaa gttttgcttg gttctactca ttactctact 600
gctgttgata tttggtctgt tggatgcac tttgccgaga tgattaggag gcaagctctt 660
ttccctgggtg attctgagtt tcagcaacta cttcatattt tcagattggt aggaacacca 720
actgagcagc aatggccggg tgtaatggca ttgcgtgact ggcattgtcta tccaaagtgg 780
gagccgcaag acttatcacg tgctgttcca tctctatctc ctgaaggaa tgatcttctc 840
acgcaaagt tgaagtacaa tccagcagaa agaatttcag caaaagcagc tctcgatcat 900
ccctactttg acagccttga caaatctcag ttctga 936

```

<210> 4
 <211> 311
 <212> PRT
 <213> Arabidopsis thaliana

<400> 4
 Met Glu Lys Tyr Glu Lys Leu Glu Lys Val Gly Glu Gly Thr Tyr Gly
 1 5 10 15
 Lys Val Tyr Lys Ala Met Glu Lys Thr Thr Gly Lys Leu Val Ala Leu
 20 25 30
 Lys Lys Thr Arg Leu Glu Met Asp Glu Glu Gly Ile Pro Pro Thr Ala
 35 40 45
 Leu Arg Glu Ile Ser Leu Leu Gln Met Leu Ser Gln Ser Ile Tyr Ile
 50 55 60
 Val Arg Leu Leu Cys Val Glu His Val Ile Gln Ser Lys Asp Ser Thr
 65 70 75 80
 Val Ser His Ser Pro Lys Ser Asn Leu Tyr Leu Val Phe Glu Tyr Leu
 85 90 95
 Asp Thr Asp Leu Lys Lys Phe Ile Asp Ser His Arg Lys Gly Ser Asn
 100 105 110
 Pro Arg Pro Leu Glu Ala Ser Leu Val Gln Arg Phe Met Phe Gln Leu
 115 120 125
 Phe Lys Gly Val Ala His Cys His Ser His Gly Val Leu His Arg Asp
 130 135 140
 Leu Lys Pro Gln Asn Leu Leu Leu Asp Lys Asp Lys Gly Ile Leu Lys
 145 150 155 160
 Ile Ala Asp Leu Gly Leu Ser Arg Ala Phe Thr Val Pro Leu Lys Ala
 165 170 175
 Tyr Thr His Glu Ile Val Thr Leu Trp Tyr Arg Ala Pro Glu Val Leu
 180 185 190
 Leu Gly Ser Thr His Tyr Ser Thr Ala Val Asp Ile Trp Ser Val Gly
 195 200 205
 Cys Ile Phe Ala Glu Met Ile Arg Arg Gln Ala Leu Phe Pro Gly Asp
 210 215 220
 Ser Glu Phe Gln Gln Leu Leu His Ile Phe Arg Leu Leu Gly Thr Pro

225											230											235											240
Thr	Glu	Gln	Gln	Trp	Pro	Gly	Val	Met	Ala	Leu	Arg	Asp	Trp	His	Val																		
				245					250								255																
Tyr	Pro	Lys	Trp	Glu	Pro	Gln	Asp	Leu	Ser	Arg	Ala	Val	Pro	Ser	Leu																		
				260					265								270																
Ser	Pro	Glu	Gly	Ile	Asp	Leu	Leu	Thr	Gln	Met	Leu	Lys	Tyr	Asn	Pro																		
				275					280								285																
Ala	Glu	Arg	Ile	Ser	Ala	Lys	Ala	Ala	Leu	Asp	His	Pro	Tyr	Phe	Asp																		
				290					295								300																
Ser	Leu	Asp	Lys	Ser	Gln	Phe																											
				305					310																								

```
<210> 5
<211> 948
<212> DNA
<213> Arabidopsis thaliana
```

[illegible]

```
<210> 6
<211> 315
<212> PRT
<213> Arabidopsis thaliana
```

```

<400>      6
Met Asp Asn Asn Gly Val Lys Pro Ala Val Ser Ala Met Glu Ala Phe
1              5              10              15

Glu Lys Leu Glu Lys Val Gly Glu Gly Thr Tyr Gly Lys Val Tyr Arg
          20              25              30

Ala Arg Glu Lys Ala Thr Gly Met Ile Val Ala Leu Lys Lys Thr Arg
          35              40              45

Leu His Glu Asp Glu Glu Gly Val Pro Pro Thr Thr Leu Arg Glu Ile
          50              55              60

Ser Ile Leu Arg Met Leu Ala Arg Asp Pro His Ile Val Arg Leu Met

```

65	70	75	80
Asp Val Lys Gln Gly Ile Asn Lys Glu Gly Lys Thr Val Leu Tyr Leu	85	90	95
Val Phe Glu Tyr Val Asp Thr Asp Leu Lys Lys Phe Ile Arg Ser Phe	100	105	110
Arg Gln Ala Gly Gln Asn Ile Pro Gln Asn Thr Val Lys Cys Leu Met	115	120	125
Tyr Gln Leu Cys Lys Gly Met Ala Phe Cys His Gly His Gly Val Leu	130	135	140
His Arg Asp Leu Lys Pro His Asn Leu Leu Met Asp Arg Lys Thr Met	145	150	155
Thr Leu Lys Ile Ala Asp Leu Gly Leu Ala Arg Ala Phe Thr Leu Pro	165	170	175
Met Lys Lys Tyr Thr His Glu Ile Leu Thr Leu Trp Tyr Arg Ala Pro	180	185	190
Glu Val Leu Leu Gly Ala Thr His Tyr Ser Thr Gly Val Asp Met Trp	195	200	205
Ser Val Gly Cys Ile Phe Ala Glu Leu Val Thr Lys Gln Ala Ile Phe	210	215	220
Ala Gly Asp Ser Glu Leu Gln Gln Leu Leu Arg Ile Phe Arg Leu Leu	225	230	235
Gly Thr Pro Asn Glu Glu Val Trp Pro Gly Val Ser Lys Leu Lys Asp	245	250	255
Trp His Glu Tyr Pro Gln Trp Lys Pro Leu Ser Leu Ser Thr Ala Val	260	265	270
Pro Asn Leu Asp Glu Ala Gly Leu Asp Leu Leu Ser Lys Met Leu Glu	275	280	285
Tyr Glu Pro Ala Lys Arg Ile Ser Ala Lys Lys Ala Met Glu His Pro	290	295	300
Tyr Phe Asp Asp Leu Pro Asp Lys Ser Ser Leu	305	310	315

```
<210> 7
<211> 1115
<212> DNA
<213> Oryza sativa
```

<400>	7						
acctctctctc	cgattaatcc	cctcccctcc	tcttctctccc	acttctgcgc	ctgctcttcc		60
tcccctcgcc	gacctacct	actcgcgccg	ccggcgctcg	attgggcggc	aaacggaggg		120
ggggttaacc	ctgatggagc	agtacgagaa	ggaggagaag	attggggagg	gcacgtacgg		180
ggtggtgtac	agggcgcggg	acaaggctcac	caacgagacg	atcgcgctca	agaagatccg		240
gcttgagcag	gaggatgagg	gcgtcccctc	caccgcaatc	cgcgagatct	cgctcctcaa		300
qtagatgcat	cacggcaaca	tcgtcaggtt	acacgatgtt	atccacagtg	agaagcgcat		360

CD103PCT.ST25.txt

```

atatacttgtc tttgagtatc tggatctgga cctaaagaag ttcattggact cttgtccaga 420
gtttgcgaaa aacccactt taattaagtc atatctctat cagatactcc gcggcggtgc 480
ttactgtcat tctcatagag ttcttcatcg agatttgaaa cctcagaatt tattgataga 540
tcggcgctact aatgcactga agcttgacga ctttggttta gccaggggcat ttggaattcc 600
tgtccgcacg tttactcacg aggttgtaac cttgtggtat agagctccag agatccttct 660
tggatcaagg cagtattcta caccagttag tatgtggtca gttggttgta tctttgcaga 720
aatgggtgaac cagaaaccac tgttccctgg tgattctgag attgatgaat tatttaagat 780
attcagggta ctaggaaactc caaatgaaca aagttggcca ggagttagct cattacctga 840
ctacaagtct gctttcccca agtggcaagc acaggatctt gcaactattg tccctactct 900
tgaccctgct ggtttggacc ttctctctaa aatgcttcgg tacgagccaa acaaaaggat 960
cacagctaga caggctcttg agcatgaata cttcaaggac cttgagatgg tacaatgacc 1020
ctgctatggc tttacattgg attggcatat gtatgggctg ggctcctcat ttcattcctt 1080
ctgtgaacgc tgtgcccttc gtttgggcat ttttg 1115

```

<210> 8
 <211> 294
 <212> PRT
 <213> Oryza sativa

<400> 8
 Met Glu Gln Tyr Glu Lys Glu Glu Lys Ile Gly Glu Gly Thr Tyr Gly
 1 5 10 15
 Val Val Tyr Arg Ala Arg Asp Lys Val Thr Asn Glu Thr Ile Ala Leu
 20 25 30
 Lys Lys Ile Arg Leu Glu Gln Glu Asp Glu Gly Val Pro Ser Thr Ala
 35 40 45
 Ile Arg Glu Ile Ser Leu Leu Lys Glu Met His His Gly Asn Ile Val
 50 55 60
 Arg Leu His Asp Val Ile His Ser Glu Lys Arg Ile Tyr Leu Val Phe
 65 70 75 80
 Glu Tyr Leu Asp Leu Asp Leu Lys Lys Phe Met Asp Ser Cys Pro Glu
 85 90 95
 Phe Ala Lys Asn Pro Thr Leu Ile Lys Ser Tyr Leu Tyr Gln Ile Leu
 100 105 110
 Arg Gly Val Ala Tyr Cys His Ser His Arg Val Leu His Arg Asp Leu
 115 120 125
 Lys Pro Gln Asn Leu Leu Ile Asp Arg Arg Thr Asn Ala Leu Lys Leu
 130 135 140
 Ala Asp Phe Gly Leu Ala Arg Ala Phe Gly Ile Pro Val Arg Thr Phe
 145 150 155 160
 Thr His Glu Val Val Thr Leu Trp Tyr Arg Ala Pro Glu Ile Leu Leu
 165 170 175
 Gly Ser Arg Gln Tyr Ser Thr Pro Val Asp Met Trp Ser Val Gly Cys
 180 185 190
 Ile Phe Ala Glu Met Val Asn Gln Lys Pro Leu Phe Pro Gly Asp Ser
 195 200 205

CD103PCT.ST25.txt

Glu Ile Asp Glu Leu Phe Lys Ile Phe Arg Val Leu Gly Thr Pro Asn
 210 215 220
 Glu Gln Ser Trp Pro Gly Val Ser Ser Leu Pro Asp Tyr Lys Ser Ala
 225 230 235 240
 Phe Pro Lys Trp Gln Ala Gln Asp Leu Ala Thr Ile Val Pro Thr Leu
 245 250 255
 Asp Pro Ala Gly Leu Asp Leu Leu Ser Lys Met Leu Arg Tyr Glu Pro
 260 265 270
 Asn Lys Arg Ile Thr Ala Arg Gln Ala Leu Glu His Glu Tyr Phe Lys
 275 280 285
 Asp Leu Glu Met Val Gln
 290
 <210> 9
 <211> 294
 <212> PRT
 <213> Oryza sativa
 <400> 9
 Met Glu Gln His Glu Lys Glu Glu Lys Ile Gly Glu Gly Thr Tyr Gly
 1 5 10 15
 Val Val Tyr Arg Ala Arg Asp Lys Val Thr Asn Glu Thr Ile Ala Leu
 20 25 30
 Lys Lys Ile Arg Leu Glu Gln Glu Asp Glu Gly Val Pro Ser Thr Ala
 35 40 45
 Ile Arg Glu Ile Ser Leu Leu Lys Glu Met His His Gly Asn Ile Val
 50 55 60
 Arg Leu His Asp Val Ile His Ser Glu Lys Arg Ile Tyr Leu Asp Phe
 65 70 75 80
 Glu Tyr Leu Asp Leu Asp Leu Lys Lys Phe Met Asp Ser Cys Pro Glu
 85 90 95
 Phe Ala Lys Asn Pro Thr Leu Ile Lys Ser Tyr Leu Tyr Gln Ile Leu
 100 105 110
 Arg Gly Val Ala Tyr Cys His Ser His Arg Val Leu His Arg Asp Leu
 115 120 125
 Lys Pro Gln Asn Leu Leu Ile Asp Arg Arg Thr Asn Ala Leu Lys Leu
 130 135 140
 Ala Asp Phe Gly Leu Ala Arg Thr Phe Gly Ile Pro Val Arg Thr Phe
 145 150 155 160
 Thr His Glu Val Val Thr Leu Trp Tyr Arg Ala Pro Glu Ile Leu Leu
 165 170 175
 Gly Ser Arg Gln Tyr Ser Thr Pro Val Asp Met Trp Ser Val Gly Cys
 180 185 190

CD103PCT.ST25.txt

```

Ile Phe Ala Glu Met Val Asn Gln Lys Pro Leu Phe Pro Gly Asp Ser
    195                200                205

Glu Ile Asp Glu Leu Phe Lys Ile Phe Arg Val Leu Gly Thr Pro Asn
    210                215                220

Glu Gln Ser Trp Pro Gly Val Ser Ser Leu Pro Asp Tyr Lys Ser Ala
    225                230                235                240

Phe Pro Lys Trp Gln Ala Gln Asp Leu Ala Thr Ile Val Pro Thr Leu
    245                250                255

Asp Pro Ala Gly Leu Asp Leu Leu Ser Lys Met Leu Arg Tyr Glu Pro
    260                265                270

Asn Lys Arg Ile Thr Ala Arg Gln Ala Leu Glu His Glu Tyr Phe Lys
    275                280                285

Asp Leu Glu Met Val Gln
    290

<210> 10
<211> 294
<212> PRT
<213> Oryza sativa

<400> 10
Met Glu Gln Tyr Glu Lys Glu Glu Lys Ile Gly Glu Gly Thr Tyr Gly
  1          5          10          15

Val Val Tyr Arg Ala Arg Asp Lys Val Thr Asn Glu Thr Thr Ala Leu
    20          25          30

Lys Lys Ile Arg Leu Glu Gln Glu Asp Glu Gly Val Pro Ser Thr Ala
    35          40          45

Ile Arg Glu Ile Ser Leu Leu Lys Glu Met His His Gly Asn Ile Val
    50          55          60

Arg Leu His Asp Val Ile His Ser Glu Lys Arg Ile Tyr Leu Val Phe
    65          70          75          80

Glu Tyr Leu Asp Leu Asp Leu Lys Lys Phe Met Asp Ser Cys Pro Glu
    85          90          95

Phe Ala Lys Asn Pro Thr Leu Ile Lys Ser Tyr Leu Tyr Gln Ile Leu
    100         105         110

Arg Gly Val Ala Tyr Cys His Ser His Arg Val Leu His Arg Asp Leu
    115         120         125

Lys Pro Gln Asn Leu Leu Ile Asp Arg Arg Thr Asn Ala Leu Lys Leu
    130         135         140

Ala Asp Phe Gly Leu Ala Arg Ala Phe Gly Ile Pro Val Arg Thr Phe
    145         150         155         160

Thr His Glu Val Val Thr Leu Trp Tyr Arg Ala Pro Glu Ile Leu Leu

```


CD103PCT.ST25.txt

```

165              170              175
Gly Ser Arg Gln Tyr Ser Thr Pro Val Asp Met Trp Ser Val Gly Cys
180              185              190
Ile Phe Ala Glu Met Val Asn Gln Lys Pro Leu Phe Pro Gly Asp Ser
195              200              205
Glu Ile Asp Glu Leu Phe Lys Ile Phe Arg Val Leu Gly Thr Pro Asn
210              215              220
Glu Gln Ser Trp Pro Gly Val Ser Ser Leu Pro Asp Tyr Lys Ser Ala
225              230              235              240
Phe Pro Lys Trp Gln Ala Gln Asp Leu Ala Thr Ile Val Pro Thr Leu
245              250              255
Asp Pro Ala Gly Leu Asp Leu Leu Ser Lys Met Leu Arg Tyr Glu Pro
260              265              270
Asn Lys Arg Ile Thr Ala Arg Gln Ala Leu Glu His Glu Tyr Phe Lys
275              280              285
Asp Leu Glu Met Val Gln
290

<210> 11
<211> 294
<212> PRT
<213> Oryza sativa

<400> 11
Met Glu Gln Tyr Val Lys Glu Glu Lys Ile Gly Glu Gly Thr Tyr Gly
1              5              10              15
Val Val Tyr Arg Ala Arg Asp Lys Val Thr Asn Glu Thr Ile Ala Leu
20              25              30
Lys Lys Ile Arg Leu Glu Gln Glu Asp Glu Gly Val Pro Ser Thr Ala
35              40              45
Ile Arg Glu Ile Ser Leu Leu Lys Glu Met His His Gly Asn Ile Val
50              55              60
Arg Leu His Asp Val Ile His Ser Glu Lys Arg Ile Tyr Leu Val Phe
65              70              75              80
Glu Tyr Leu Asp Leu Asp Leu Lys Lys Phe Met Asp Ser Cys Pro Glu
85              90              95
Phe Ala Lys Asn Pro Thr Leu Ile Lys Ser Tyr Leu Tyr Gln Ile Leu
100             105             110
Arg Gly Val Ala Tyr Cys His Ser His Ser Val Leu His Arg Asp Leu
115             120             125
Lys Pro Gln Asn Leu Leu Ile Asp Arg Arg Thr Asn Ala Leu Glu Leu
130             135             140

```

CD103PCT.ST25.txt

Ala Asp Phe Gly Leu Ala Arg Ala Phe Gly Ile Pro Val Arg Thr Phe
 145 150 155 160
 Thr His Glu Val Val Thr Leu Trp Tyr Arg Ala Pro Glu Ile Leu Leu
 165 170 175
 Gly Ser Arg Gln Tyr Ser Thr Pro Val Asp Met Trp Ser Val Gly Cys
 180 185 190
 Ile Phe Ala Glu Met Val Asn Gln Lys Pro Leu Phe Pro Gly Asp Ser
 195 200 205
 Glu Ile Asp Glu Leu Phe Lys Ile Phe Arg Val Leu Gly Thr Pro Asn
 210 215 220
 Glu Gln Ser Trp Pro Gly Val Ser Ser Leu Pro Asp Tyr Lys Ser Ala
 225 230 235 240
 Phe Pro Lys Trp Gln Ala Gln Asp Leu Ala Thr Ile Val Pro Thr Leu
 245 250 255
 Asp Pro Ala Gly Leu Asp Leu Leu Ser Lys Met Leu Arg Tyr Glu Pro
 260 265 270
 Asn Lys Arg Ile Thr Ala Arg Gln Ala Leu Glu His Glu Tyr Phe Lys
 275 280 285
 Asp Leu Glu Met Val Gln
 290
 <210> 12
 <211> 294
 <212> PRT
 <213> Oryza sativa
 <400> 12
 Met Glu Gln Tyr Glu Lys Glu Glu Lys Ile Gly Glu Gly Thr Tyr Gly
 1 5 10 15
 Val Val Tyr Arg Ala Arg Asp Lys Val Thr Asn Glu Thr Ile Ala Leu
 20 25 30
 Lys Lys Ile Arg Leu Glu Gln Glu Asp Glu Gly Val Pro Ser Thr Ala
 35 40 45
 Ile Arg Glu Ile Ser Leu Leu Lys Glu Met His His Gly Asn Ile Val
 50 55 60
 Arg Leu His Asp Val Ile His Ser Glu Lys Arg Ile Tyr Leu Val Phe
 65 70 75 80
 Glu Tyr Leu Asp Leu Asp Leu Lys Lys Phe Met Asp Ser Cys Pro Glu
 85 90 95
 Phe Ala Lys Asn Pro Thr Leu Ile Lys Ser Tyr Leu Tyr Gln Ile Leu
 100 105 110
 Arg Gly Val Ala Tyr Cys His Ser His Arg Val Leu His Arg Asp Leu
 115 120 125

CD103PCT.ST25.txt

Lys Pro Gln Asn Leu Leu Ile Asp Arg Arg Thr Asn Ala Leu Lys Leu
 130 135 140
 Ala Asp Phe Gly Leu Ala Arg Ala Phe Arg Ile Pro Val Arg Thr Phe
 145 150 155 160
 Thr His Glu Val Val Thr Leu Trp Tyr Arg Ala Pro Glu Ile Leu Leu
 165 170 175
 Gly Ser Arg Gln Tyr Ser Thr Pro Val Asp Met Trp Ser Val Gly Cys
 180 185 190
 Ile Phe Ala Glu Met Val Asn Gln Lys Pro Leu Phe Pro Gly Asp Ser
 195 200 205
 Glu Ile Asp Glu Leu Phe Lys Ile Phe Arg Val Leu Gly Thr Pro Asn
 210 215 220
 Glu Gln Ser Trp Pro Gly Val Ser Ser Leu Pro Asp Tyr Lys Ser Ala
 225 230 235 240
 Phe Pro Lys Trp Gln Ala Gln Asp Leu Ala Thr Ile Val Pro Thr Leu
 245 250 255
 Asp Pro Ala Gly Leu Asp Leu Leu Ser Lys Met Leu Arg Tyr Glu Pro
 260 265 270
 Asn Lys Arg Ile Thr Ala Arg Gln Ala Leu Glu His Glu Tyr Phe Lys
 275 280 285
 Asp Leu Glu Met Val Gln
 290
 <210> 13
 <211> 294
 <212> PRT
 <213> Oryza sativa
 <400> 13
 Met Glu Pro Tyr Glu Lys Glu Glu Lys Ile Gly Glu Gly Thr Tyr Gly
 1 5 10 15
 Val Val Tyr Arg Ala Arg Asp Lys Val Thr Asn Glu Thr Ile Ala Leu
 20 25 30
 Lys Lys Ile Arg Leu Ala Gln Glu Asp Glu Gly Val Pro Ser Thr Ala
 35 40 45
 Ile Arg Glu Ile Ser Leu Leu Lys Glu Met His His Gly Asn Ile Val
 50 55 60
 Arg Leu His Asp Val Ile His Ser Glu Lys Arg Ile Tyr Leu Val Phe
 65 70 75 80
 Glu Tyr Leu Asp Leu Asp Leu Lys Lys Phe Met Asp Ser Cys Pro Glu
 85 90 95
 Phe Ala Lys Asn Pro Thr Leu Ile Lys Ser Tyr Leu Tyr Gln Ile Leu

CD103PCT.ST25.txt

100 105 110
 Arg Gly Val Ala Tyr Cys His Ser His Arg Val Leu His Arg Asp Leu
 115 120 125
 Lys Pro Gln Asn Leu Leu Ile Asp Leu Arg Thr Asn Ala Leu Lys Leu
 130 135 140
 Ala Asp Phe Gly Leu Ala Arg Ala Phe Gly Ile Pro Val Arg Thr Phe
 145 150 155 160
 Thr His Glu Val Val Thr Leu Trp Tyr Arg Ala Pro Glu Ile Leu Leu
 165 170 175
 Gly Ser Arg Gln Tyr Ala Thr Pro Val Asp Met Trp Ser Val Gly Cys
 180 185 190
 Thr Phe Ala Glu Met Val Asn Gln Lys Pro Leu Phe Pro Gly Asp Ser
 195 200 205
 Glu Ile Asp Glu Leu Phe Lys Ile Phe Arg Val Leu Gly Thr Pro Asn
 210 215 220
 Glu Gln Ser Trp Pro Gly Val Ser Ser Leu Pro Asp Tyr Lys Ser Ala
 225 230 235 240
 Phe Pro Lys Trp Gln Ala Gln Asp Leu Ala Thr Ile Val Pro Thr Leu
 245 250 255
 Asp Pro Ala Gly Leu Asp Leu Leu Ser Lys Val Leu Arg Tyr Glu Pro
 260 265 270
 Asn Lys Arg Ile Thr Ala Gln Gln Ala Leu Glu His Glu Tyr Phe Lys
 275 280 285
 Asp Leu Glu Met Val Gln
 290

<210> 14

<211> 1243

<212> DNA

<213> Oryza sativa

<400> 14

aaaaccaccg	agggacctga	tctgcaccgg	ttttgatagt	tgagggaccc	gttggtgtctg	60
gttttccgat	cgagggacga	aaatcggatt	cggtgtaaag	ttaagggacc	tcagatgaac	120
ttattccgga	gcatgattgg	gaagggagga	cataaggccc	atgtcgcatg	tgtttggacg	180
gtccagatct	ccagatcact	cagcaggatc	ggccgcgttc	gcgtagcacc	cgcggtttga	240
ttcggcttcc	cgcaaggcgg	cggccgggtg	ccgtgccgcc	gtagcttccg	ccggaagcga	300
gcacgccgcc	gccgccgacc	cggtctctgc	tttgcaccgc	cttgcacgcg	atacatcggg	360
atagatagct	actactctct	cgttttcaca	atgtaaatac	ttctactatt	ttccacattc	420
atattgatgt	taatgaatat	agacatatat	atctatttag	attcattaac	atcaatatga	480
atgtaggaaa	tgctagaatg	acttacattg	tgaattgtga	aatggacgaa	gtacctacga	540
tggtatggatg	caggatcatg	aaagaattaa	tgcaagatcg	tatctgccgc	atgcaaaatc	600
ttactaatgt	cgctgcatat	atgcatgaca	gcctgcatgc	ggcggtgtaa	gcgtgttcac	660
ccattaggaa	gtaaccttgt	cattacttat	accagtacta	catactatat	agtattgatt	720
tcatgagcaa	atctacaaaa	ctggaaagca	ataagaaata	cgggactgga	aaagactcaa	780
cattaatcac	caaatatttc	gccttctcca	gcagaatata	tatctctcca	tcttgatcac	840
tgtacacact	gacagtgtac	gcataaacgc	agcagccagc	ttaactgtcg	tctcaccgct	900

CD103PCT.ST25.txt

```

gcacactggc cttccatctc aggctagctt tctcagccac ccacgtgaca tgtcaactcg      960
gcgcgcgcac aggcacaaat tacgtacaaa acgcatgacc aaatcaaaac caccggagaa      1020
gaatcgctcc cgcgcgcggc ggcgacgcgc acgtacgaac gcacgcacgc acgcccaccc      1080
ccacgcacag atcgcgcgcg acgcccgcga caccggcgcg ccacccgcgc cctcacctcg      1140
ccgactataa atacgtaggc atctgcttga tcttgtcatc catctacca ccaaaaaaaaa      1200
aaggaaaaaa aaacaaaaca caccaagcca aataaaagcg aca                        1243

```

<210> 15
 <211> 2191
 <212> DNA
 <213> *Oryza sativa*

```

<400> 15
aatccgaaaa gtttctgcac cgttttcacc ccctaactaa caatataggg aacgtgtgct      60
aatataaaaa tgagacctta tataatgtagc gctgataact agaactatgc aagaaaaact      120
catccaccta ctttagtggc aatcgggcta aataaaaaag agtcgctaca ctagtttcgt      180
tttcttagt aattaagtgg gaaaatgaaa tcattattgc ttagaatata cgttcacatc      240
tctgtcatga agttaaatta ttcgaggtag ccataattgt catcaaacctc ttcttgaata      300
aaaaaatctt tctagctgaa ctcaatgggt aaagagagag atttttttta aaaaaataga      360
atgaagatat tctgaacgta ttggcaaaga tttaaacata taattatata attttatagt      420
ttgtgcattc gtcatatcgc acatcattaa ggacatgtct tactccatcc caatttttat      480
ttagtaatta aagacaattg acttattttt attatttacc ttttttcgat tagatgcaag      540
gtacttacgc acacactttg tgctcatgtg catgtgtgag tgcacctcct caatacacgt      600
tcaactagca acacatctct aatatcactc gcctatttaa tacatttagg tagcaatatc      660
tgaattcaag cactccacca tcaccagacc acttttaata atatctaaaa tacaataaat      720
aattttacag aatagcatga aaagtatgaa acgaactatt taggtttttc acatacaaaa      780
aaaaaaagaa ttttgctcgt gcgcgagcgc caatctccca tattgggcac acaggcaaca      840
acagagtggc tgcccacaga acaaccaca aaaaacgatg atctaacgga ggacagcaag      900
tccgcaacaa ccttttaaca gcaggctttg cgccaggag agaggaggag agggcaagaa      960
aaccaagcat cctcctcctc ccatctataa attcctcccc ccttttcccc tctctatata      1020
ggaggcatcc aagccaagaa gagggagagc accaaggaca cgcgactagc agaagccgag      1080
cgaccgcctt cttcgatcca tatcttcggt tcgagttctt ggtcgatctc ttccctcctc      1140
cacctcctcc tcacagggtg tgtgcccctc ggttgttctt ggatttattg ttctagggtg      1200
tgtagtacgg gcgttgatgt taggaaaggg gatctgtatc tgtgatgatt cctgttcttg      1260
gatttgggat agaggggttc ttgatgttgc atgttatcgg ttcggtttga ttagtagtat      1320
ggttttcaat cgtctggaga gctctatgga aatgaaatgg tttagggtac ggaatcttgc      1380
gattttgtga taccttttgt ttgaggtaaa atcagagcac cgggtgattt gcttggtgta      1440
ataaaagtac ggttgtttgg tcctcgattc tggtagtgat gcttctcgat ttgacgaagc      1500
tatcctttgt ttattcccta ttgaacaaaa ataattcaac tttgaagacg gtcccgttga      1560
tgagattgaa tgattgattc ttaagcctgt ccaaaatttc gcagctggct tgtttagata      1620
cagtagtccc catcacgaaa ttcatggaaa cagttataat cctcaggaaac aggggattcc      1680
ctgttcttcc gatttgcctt agtcccagaa ttttttttcc caaatatctt aaaaagtcac      1740
tttctgggtc agttcaatga attgattgct acaataatg cttttatagc gttatcctag      1800
ctgtagttca gttaataggt aatacccta tagtttagtc aggagaagaa cttatccgat      1860
ttctgatctc catttttaat tatatgaaat gaactgtagc ataagcagta ttcatctgga      1920
ttattttttt tttagctctc accccttcat tattctgagc tgaaagtctg gcatgaactg      1980
tctcaattt tgttttcaaa ttcacatcga ttatctatgc attatcctct tgtatctacc      2040
tgtagaagtt tctttttggg tattccttga ctgcttgatt acagaaagaa atttatgaag      2100
ctgtaatcgg gatagttata ctgcttgctc ttatgattca tttcctttgt gcagttcttg      2160
gtgtagcttg ccactttcac cagcaaagtt c

```

<210> 16
 <211> 57
 <212> DNA
 <213> Artificial sequence

<220>
 <223> sense primer: prm0350

CD103PCT.ST25.txt

<400> 16
ggggacaagt ttgtacaaaa aagcaggctt cacaatggag aagtacgaga agctaga 57

<210> 17
<211> 51
<212> DNA
<213> Artificial sequence

<220>
<223> antisense primer: prm0351

<400> 17
ggggaccact ttgtacaaga aagctggggtt cagaactgag acttgtcaag g 51

<210> 18
<211> 55
<212> DNA
<213> Artificial sequence

<220>
<223> sense primer: prm439

<400> 18
ggggacaagt ttgtacaaaa aagcaggctt cacaatggag aaatacgaga agctc 55

<210> 19
<211> 49
<212> DNA
<213> Artificial sequence

<220>
<223> antisense primer: prm440

<400> 19
ggggaccact ttgtacaaga aagctgggtg gtcagaactg agatttgtc 49

<210> 20
<211> 54
<212> DNA
<213> Artificial sequence

<220>
<223> sense primer: prm2213

<400> 20
ggggacaagt ttgtacaaaa aagcaggctt cacaatggac aacaatggag ttaa 54

<210> 21
<211> 49
<212> DNA
<213> Artificial sequence

<220>
<223> antisense primer: prm2214

<400> 21
ggggaccact ttgtacaaga aagctggggtt cagagagagg acttgtcag 49